

The rapid on-line diffusion of short messages, or “memes,” can be mathematically treated as an epidemic spreading process. Introduction The spread of information in on-line social systems bears striking resemblance to pathogen transmission in populations. Classic well-mixed models offer a parsimonious answer: inoculating a fraction  $1 - 1/R_0$  of individuals collapses the basic reproduction number  $R_0$  to zero.

In this study we revisit the archetypal configuration model under the simplifying assumption that random vaccination, each node is independently immunized with probability  $p$ .

**Random vaccination**—each node is independently immunized with probability  $p$ .  
**Fixed degree vaccination**—all nodes of degree exactly  $k = 10$  are vaccinated.

The remainder of the paper is organized as follows. Section 3 details the analysis of the

The remainder of the paper is organized as follows. Section 2 details the analytical framework, network construction, parameter estimation, and model selection.

Methodology Analytical vaccination threshold on uncorrelated networks For configuration networks without degree corr

When immunization is *not* random but concentrated on a specific degree  $k$ , the algebra changes. Let  $P_k$  be the pre-vaccination probability that a randomly chosen node has degree  $k$ . Then  $\langle k^2 \rangle' = \frac{\langle k^2 \rangle - (k)^2 f}{1-f}$ . Plugging these into  $q'$  yields

Selecting a representative degree distribution To validate the analytics in silico we require a concrete network whose me

so vaccinating all  $k = 10$  nodes removes only 0.81% of the population.

Network synthesis Using the `networkx` configuration-model generator, we sampled  $N = 10^4$  degrees from the selected model.

$q = 3.9623$  ( $\approx 4$ ). The adjacency matrix was stored as a CSR matrix (`network.npz`) for simulation reproducibility.

Disease dynamics and parameters The memo dynamics were modeled as a continuous-time SIR process with infection rate

Vaccination scenarios and initialization **Scenario 1 – Random vaccination.** Each node was independently assigned

**Scenario 2 – Fixed-degree vaccination.** All nodes of degree exactly 10 were pre-assigned to  $R$ . This immunized fraction of the population was constant at approximately 10%.

Simulation engine We employed **fastGEMF** v2.1, which implements the generalized epidemic mean-field framework via the Gillespie algorithm.

Results Analytical predictions Equation eq:rand\_threshold\_prescribes\_pc = 0.75 for random immunization. For fixed-degr

Stochastic simulations validate analytics \*Scenario 1 (random 75% vaccination). The epidemic never gained traction (Figure 1).

\*Scenario 2 (all  $k = 10$  vaccinated). Figure shows a dramatic outbreak: prevalence climbed to 15.9% around  $t = 23$  an-

<http://i111.8.21.128/~T...> Temporal evolution of compartments under 75% random vaccination. The infectious cur-

[http://width=0.9]results-12.png Temporal evolution under universal degree-10 vaccination. A large-scale outbreak occurs immediately after the initial vaccination.

**Discussion** The combined analytical-computational investigation yields three principal insights. Random versus targeted immunization efficiency. Random vaccination achieves herd immunity exactly at the classical basic reproduction number.

Random versus targeted immunization efficiency. Random vaccination achieves herd immunity exactly at the classical b<sub>c</sub>. Implications for social media content moderation. Online platforms often rely on flagging or throttling a handful of large

Implications for social media content moderation. Online platforms often rely on flagging or filtering a handful of large, static contact structures, and no degree-degree correlation

**Conclusion** We analytically derived and numerically validated vaccination thresholds to halt meme propagation on an individual level.

Conclusion We analytically derived and numerically validated vaccination thresholds to halt disease propagation on an SIR graph.

M. Zhou, W. M. Xiong, H. Liao, *et al.*, "Analytical connection between thresholds and immunization strategies of SIS model," *Journal of Mathematical Analysis and Applications*, vol. 405, no. 2, pp. 529–541, 2013.

J. B. Jia, W. Shi, P. Yang, et al., "Immunization strategies in directed networks," *Math. Biosci. Eng.*, vol. 17, no. 4, pp. 39

Reproducibility material All Python scripts, networks, and CSV outputs are included in the accompanying `output` directory.